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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2008; month=8; day=1; hr=17; min=1; sec=20; ms=803; ]

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Reviewer Comments:

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"Artificial Peptide Sequence", "Artificial amino acid sequence", and "Artificial nucleic acid sequence" are all insufficient responses for numeric identifier <223>. Please explain the source of the genetic material. If the sequence is put together from several organisms, please list those organisms. If the sequence is made in the laboratory, please

indicate that the sequence is synthesized. These errors appear in other sequences in the sequence listing. Please make all necessary changes.

\*\*\*\*\*

Application No: 09631613 Version No: 2.0

**Input Set:****Output Set:**

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**Finished:** 2008-06-27 14:14:19.635  
**Elapsed:** 0 hr(s) 0 min(s) 2 sec(s) 356 ms  
**Total Warnings:** 83  
**Total Errors:** 0  
**No. of SeqIDs Defined:** 89  
**Actual SeqID Count:** 89

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**Input Set:**

**Output Set:**

**Started:** 2008-06-27 14:14:17.279  
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Error code

Error Description

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W 402

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Hansen, Connie J

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<150> US 08/957,709

<151> 1997-10-24

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ccttatgcat ggaatttgcc cacgggaaat ccagtcataa ctgagatcac tggatttatc      240

gagcatgttg agttagcagg ggaacatgag aataaagcag atttaatttt ggtttgcct      300

gccactgcca acacaattag taagattgca tgtggaatag atgatactcc agtaactaca      360

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tacagagtta ttaaaaacgt ccacaaaaaa acattggaag ggaagagagt cctagtaacg      600

gcgggagcaa caagagagta catagatcca ataagattca taacaaatgc cagcagtgga      660

aaaatgggag tagcgttggc tgaagaagca gatttttagag gagctgttac cctcataaga      720

acaaagggaa gtgtaaaggc ttttagaatc agaaaaatca aattgaaggt tgagacagtg      780

gaagaaatgc tttcagcgat tgaaaatgag ttgaggagta aaaagtatga cgtagttatt      840

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35 40 45

Ala Val Met Ser Glu Ala Ala Thr Lys Ile Ile His Pro Tyr Ala Trp  
50 55 60

Asn Leu Pro Thr Gly Asn Pro Val Ile Thr Glu Ile Thr Gly Phe Ile  
65 70 75 80

Glu His Val Glu Leu Ala Gly Glu His Glu Asn Lys Ala Asp Leu Ile  
85 90 95

Leu Val Cys Pro Ala Thr Ala Asn Thr Ile Ser Lys Ile Ala Cys Gly  
100 105 110

Ile Asp Asp Thr Pro Val Thr Thr Val Val Thr Thr Ala Phe Pro His  
115 120 125

Ile Pro Ile Met Ile Ala Pro Ala Met His Glu Thr Met Tyr Arg His  
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Pro Ile Val Arg Glu Asn Ile Glu Arg Leu Lys Lys Leu Gly Val Glu  
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Phe Ile Gly Pro Arg Ile Glu Glu Gly Arg Ala Lys Val Ala Ser Ile

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Glu Gly Lys Arg Val Leu Val Thr Ala Gly Ala Thr Arg Glu Tyr Ile		
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225	230	235 240
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260	265	270
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Arg Pro Lys Ile Lys Ala Glu Gly Lys Ile Lys Ser Gly Arg Ser Ile		
290	295	300
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